SEQUENCE LISTING

<110> RUSCH, Douglas et al

<120> ISOLATED HUMAN KINASE PROTEINS, NUCLEIC
ACID MOLECULES ENCODING HUMAN KINASE PROTEINS, AND USES
THEREOF

<130> CL001099-CIP-DIV2

<140> To be assigned
<141> 2003-10-31

<140> 10/274,878
<141> 2002-10-22

<150> 09/849,334
<151> 2001-05-07

<150> 09/773,371
<151> 2001-02-01

<160> 4

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<212> DNA
<213> Homo sapiens

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<211> 723
<212> PRT
<213> Homo sapiens

<400> 2
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        35                  40                  45
Gly Cys Arg Val His Arg Gly Glu Asp Pro Val Arg Val His Val Gly
    50                  55                  60
Pro Met Asn Pro Gln Leu His Ala Val Gly Cys Asp Ser Leu Thr Gln
65                  70                  75                  80
Ile Gln Cys Gly Gln Leu Gln Ser Arg Arg Ala Gln Ile His Gln Gln
                85                  90                  95
Ile Asp Lys Glu Leu Gln Met Arg Thr Gly Ala Glu Asn Leu Tyr Arg
            100                 105                 110
Ala Thr Ser Asn Asn Arg Val Arg Glu Thr Val Ala Leu Glu Leu Ser
            115                 120                 125
Tyr Val Asn Ser Asn Leu Gln Leu Leu Lys Glu Glu Leu Glu Glu Leu
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Ser Gly Gly Val Asp Pro Gly Arg His Gly Ser Glu Ala Val Thr Val
145                 150                 155                 160
Pro Met Ile Pro Leu Gly Leu Lys Glu Thr Lys Glu Leu Asp Trp Ser
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Thr Pro Leu Lys Glu Leu Ile Ser Val His Phe Gly Glu Asp Gly Ala
            180                 185                 190
Ser Tyr Glu Ala Glu Ile Arg Glu Leu Glu Ala Leu Arg Gln Ala Met
            195                 200                 205
Arg Thr Pro Ser Arg Asn Glu Ser Gly Leu Glu Leu Leu Thr Ala Tyr
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Tyr Asn Gln Leu Cys Phe Leu Asp Ala Arg Phe Leu Thr Pro Ala Arg
225                 230                 235                 240
Ser Leu Gly Leu Phe Phe His Trp Tyr Asp Ser Leu Thr Gly Val Pro
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2

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Thr Glu Gly Ala Arg Arg Ala Met Glu Ala Phe Gln Arg Ala Ala Gly
        290                 295                 300
Ala Phe Ser Leu Leu Arg Glu Asn Phe Ser His Ala Pro Ser Pro Asp
305                 310                 315                 320
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                325                 330                 335
Ala Gln Glu Cys Val Phe Glu Gly Leu Ser Pro Pro Ala Ser Met Ala
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Pro Gln Asp Cys Leu Ala Gln Leu Arg Leu Ala Gln Glu Ala Ala Gln
        355                 360                 365
Val Ala Ala Glu Tyr Arg Leu Val His Arg Thr Met Ala Gln Pro Pro
    370                 375                 380
Val His Asp Tyr Val Pro Val Ser Trp Thr Ala Leu Val His Val Lys
385                 390                 395                 400
Ala Glu Tyr Phe Arg Ser Leu Ala His Tyr His Val Ala Met Ala Leu
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Cys Asp Gly Ser Pro Ala Thr Glu Gly Glu Leu Pro Thr His Glu Gln
                420                 425                 430
Val Phe Leu Gln Pro Pro Thr Ser Ser Lys Pro Arg Gly Pro Val Leu
                435                 440                 445
Pro Gln Glu Leu Glu Glu Arg Arg Gln Leu Gly Lys Ala His Leu Lys
        450                 455                 460
Arg Ala Ile Leu Gly Gln Glu Glu Ala Leu Arg Leu His Ala Leu Cys
465                 470                 475                 480
Arg Val Leu Arg Glu Val Asp Leu Leu Arg Ala Val Ile Ser Gln Thr
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Leu Gln Arg Ser Leu Ala Lys Tyr Ala Glu Leu Asp Arg Glu Asp Asp
                500                 505                 510
Phe Cys Glu Ala Ala Glu Ala Pro Asp Ile Gln Pro Lys Thr His Gln
        515                 520                 525
Lys Pro Glu Ala Arg Met Pro Arg Leu Ser Gln Gly Lys Gly Pro Asp
        530                 535                 540
Ile Phe His Arg Leu Gly Pro Leu Ser Val Phe Ser Ala Lys Asn Arg
545                 550                 555                 560
Trp Arg Leu Val Gly Pro Val His Leu Thr Arg Gly Glu Gly Gly Phe
                565                 570                 575
Gly Leu Thr Leu Arg Gly Asp Ser Pro Val Leu Ile Ala Ala Val Ile
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Pro Gly Ser Gln Ala Ala Ala Gly Leu Lys Glu Gly Asp Tyr Ile
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Val Ser Val Asn Gly Gln Pro Cys Arg Trp Trp Arg His Ala Glu Val
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Val Thr Glu Leu Lys Ala Ala Gly Glu Ala Gly Ala Ser Leu Gln Val
625                 630                 635                 640
Val Ser Leu Leu Pro Ser Ser Arg Leu Pro Ser Leu Gly Asp Arg Arg
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Pro Val Leu Leu Gly Pro Arg Gly Leu Leu Arg Ser Gln Arg Glu His
            660                 665                 670
Gly Cys Lys Thr Pro Ala Ser Thr Trp Ala Ser Pro Arg Pro Leu Leu
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3

Gly Trp Pro

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<211> 19025
<212> DNA
<213> Homo sapiens

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Val Gln Asn Gln Pro Gly Gln Leu Gln Ser His Arg Ala Arg Leu His
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9

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Glu Leu Ser Thr Ser Val Asp Val Asp Gln Pro Glu Gly Glu Gly Ile
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Thr Ile Pro Met Ile Pro Leu Gly Leu Lys Glu Thr Lys Glu Leu Asp
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